ENTERED

RAW SEQUENCE LISTING   DATE: 08/04/2003
PATENT APPLICATION: **US/09/963,693A**   TIME: 14:43:17

Input Set : **N:\Crf3\RULE60\09963693.raw.txt**
Output Set: **N:\CRF4\08042003\I963693A.raw**

```
  4 <110> APPLICANT: Ruvkun, Gary
  5        Ogg, Scott
  8 <120> TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
  9        IMPAIRED GLUCOSE TOLERANCE CONDITIONS
 12 <130> FILE REFERENCE: 00786/351004
 14 <140> CURRENT APPLICATION NUMBER: 09/963,693A
 15 <141> CURRENT FILING DATE: 2001-09-25
 17 <150> PRIOR APPLICATION NUMBER: 09/205,658
 18 <151> PRIOR FILING DATE: 1998-12-03
 20 <150> PRIOR APPLICATION NUMBER: 08/857,076
 21 <151> PRIOR FILING DATE: 1997-05-15
 23 <150> PRIOR APPLICATION NUMBER: 08/888,534
 24 <151> PRIOR FILING DATE: 1997-07-07
 26 <150> PRIOR APPLICATION NUMBER: US98/10080
 27 <151> PRIOR FILING DATE: 1998-05-15
 29 <160> NUMBER OF SEQ ID NOS: 328
 31 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 33 <210> SEQ ID NO: 1
 34 <211> LENGTH: 20
 35 <212> TYPE: DNA
 36 <213> ORGANISM: Artificial Sequence
 38 <220> FEATURE:
 39 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
 41 <400> SEQUENCE: 1
 42 cgctacggca aaaaagtgaa                                          20
 44 <210> SEQ ID NO: 2
 45 <211> LENGTH: 18
 46 <212> TYPE: DNA
 47 <213> ORGANISM: Artificial Sequence
 49 <220> FEATURE:
 50 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
 52 <400> SEQUENCE: 2
 53 cgatgatgaa gatacccc                                           18
 55 <210> SEQ ID NO: 3
 56 <211> LENGTH: 20
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Artificial Sequence
 60 <220> FEATURE:
 61 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
 63 <400> SEQUENCE: 3
 64 tgatgcgaac ggcgatcgat                                         20
 66 <210> SEQ ID NO: 4
 67 <211> LENGTH: 20
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 68 <212> TYPE: DNA
 69 <213> ORGANISM: Artificial Sequence
 71 <220> FEATURE:
 72 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
 74 <400> SEQUENCE: 4
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 77 <210> SEQ ID NO: 5
 78 <211> LENGTH: 22
 79 <212> TYPE: DNA
 80 <213> ORGANISM: Artificial Sequence
 82 <220> FEATURE:
 83 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
 85 <400> SEQUENCE: 5
 86 ggtttaatta cccaagtttg ag                                       22
 88 <210> SEQ ID NO: 6
 89 <211> LENGTH: 20
 90 <212> TYPE: DNA
 91 <213> ORGANISM: Artificial Sequence
 93 <220> FEATURE:
 94 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
 96 <400> SEQUENCE: 6
 97 gctcacgggt cacacaacga                                          20
 99 <210> SEQ ID NO: 7
100 <211> LENGTH: 20
101 <212> TYPE: DNA
102 <213> ORGANISM: Artificial Sequence
104 <220> FEATURE:
105 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
107 <400> SEQUENCE: 7
108 tgatgcgaac ggcgatcgat                                          20
110 <210> SEQ ID NO: 8
111 <211> LENGTH: 21
112 <212> TYPE: DNA
113 <213> ORGANISM: Artificial Sequence
115 <220> FEATURE:
116 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
118 <400> SEQUENCE: 8
119 tgagggccaa ctaaagaaga c                                        21
121 <210> SEQ ID NO: 9
122 <211> LENGTH: 20
123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
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130 cgctacggca aaaaagtgaa                                          20
132 <210> SEQ ID NO: 10
133 <211> LENGTH: 20
134 <212> TYPE: DNA
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137 <220> FEATURE:
138 <223> OTHER INFORMATION: Primer/probe derived from C. elegans
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143 <210> SEQ ID NO: 11
144 <211> LENGTH: 5816
145 <212> TYPE: DNA
146 <213> ORGANISM: Caenorhabditis elegans
148 <220> FEATURE:
149 <221> NAME/KEY: misc_feature
150 <222> LOCATION: (1)...(5816)
151 <223> OTHER INFORMATION: n = A,T,C or G
153 <400> SEQUENCE: 11
154 ggtttaatta cccaagtttg agctccaaga gcacacatct gatcgtcgga ttctactgta    60
155 ctccccgaaa aaccaacaaa aaacacaagt ttttgaacac ttgtaaatgc agacagaacg   120
156 atgacgagaa tgaatattgt cagatgtcgg agacgacaca aaattttgga aaatttggaa   180
157 gaagagaatc tcggcccgag ctgctcgtcg acgacttcaa caaccgctgc caccgaagct   240
158 ctcggaacaa ccactgagga tatgaggctt aagcagcagc gaagctcgtc gcgtgccacg   300
159 gagcacgata ttgtcgacgg caatcaccac gacgacgagc acatcacaat gagacggctt   360
160 cgacttgtca aaaattcgcg gacgcggcgt agaacgacgc ccgattcaag tatggactgc   420
161 tatgaggaaa acccgccatc acaaaaactt caataaatta ttcttggatt tctaaaaagt   480
162 catcaatgac gtcattaatg cttttactgc tattcgcttt tgtacagccg tgtgcctcaa   540
163 tagtcgaaaa acgatgcggc ccaatcgata ttcgaaatag gccgtgggat attaagccgc   600
164 aatggtcgaa acttggtgat ccgaacgaaa aagatttggc tggtcagaga atggtcaact   660
165 gcacagtggt ggaaggttcg ctgacaatct catttgtact gaaacacaag acaaaagcac   720
166 aagaagaaat gcatcgaagt ctacagccaa gatattccca agacgaattt atcacttttc   780
167 cgcatctacg tgaaattact ggaactctgc tcgttttttga gactgaagga ttagtggatt   840
168 tgcgtaaaat tttcccaaat cttcgtgtaa ttggaggccg ttcgctgatt caacactatg   900
169 cgctgataat ttatcgaaat ccggatttgg agatcggtct tgacaagctt tccgtaattc   960
170 gaaatggtgg tgtacggata atcgataatc gaaaactgtg ctacacgaaa acgattgatt  1020
171 ggaaacattt gatcacttct tccatcaacg atgttgtcgt tgataatgct gccgagtacg  1080
172 ctgtcactga gactggattg atgtgcccac gtggagcttg cgaagaggat aaaggcgaat  1140
173 caaagtgtca ttatttggag gaaaagaatc aggaacaagg tgtcgaaaga gttcagagtt  1200
174 gttggtcgaa caccacttgc caaaagtctt gtgcttatga tcgtcttctt ccaacgaaag  1260
175 aaatcggacc gggatgtgat gcgaacggcg atcgatgtca cgatcaatgc gtgggcggtt  1320
176 gtgagcgtgt gaatgatgcc acagcatgcc acgcgtgcaa gaatgtctat cacaagggaa  1380
177 agtgtatcga aaagtgtgat gctcacctgt accttctcct tcaacgtcgt tgtgtgaccc  1440
178 gtgagcagtg tctgcagctg aatccggtgc tctcgaacaa aacagtgcct atcaaggcga  1500
179 cggcaggcct ttgctcggat aaatgtcccg atggttatca aatcaacccg gatgatcatc  1560
180 gagaatgccg aaaatgcgtt ggcaagtgtg agattgtgtg cgagatcaat cacgtcattg  1620
181 atacgtttcc gaaggcacag gcgatcaggc tatgcaatat tattgacgga aatctgacga  1680
182 tcgagattcg cggaaaacag gattcgggaa tggcgtccga gttgaaggat atatttgcga  1740
183 acattcacac gatcaccggc tacctgttgg tacgtcaatc gtcaccgttt atctcgttga  1800
184 acatgttccg gaatttacga cgtattgagg caaagtcact gttcagaaat ctatatgcta  1860
185 tcacagtttt tgaaaatccg aatttaaaaa agctattcga ttcaacgacg gatttgacgc  1920
186 ttgatcgtgg aactgtgtca attgccaata acaagatgtt atgcttcaag tatatcaagc  1980
187 agctaatgtc aaagttaaat ataccactcg atccgataga tcaatcagaa gggacaaatg  2040
W--> 188 gtgagaaggn aatctgtgag gatatggcaa tcaacgtgag catcacagcg gtcaacgcgg  2100
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189 actcggtctt ctttagttgg ccctcattca acattaccga tatagatcag cgaaagtttc 2160
190 tcggctacga gctcttcttc aaagaagtcc cacgaatcga tgagaacatg acgatcgaag 2220
191 aggatcgaag tgcgtgtgtc gattcgtggc agagtgtctt caaacagtac tacgagacgt 2280
192 cgaacggtga accgaccccg gacatttta tggatattgg accgcgcgag cgaattcggc 2340
193 cgaatacgct ctacgcgtac tatgtggcga cgcagatggt gttgcatgcc ggtgcgaaga 2400
194 acggtgtatc gaagattggt tttgtgagga cgagctacta tacgcctgat cctccgacgt 2460
195 tggcactagc gcaagtcgat tcggacgcta ttcatattac gtgggaagcg ccgctccaac 2520
196 cgaacggaga cctcacgcat tacacaatta tgtggcgtga gaatgaagtg agcccgtacg 2580
197 aggaagccga aaagttttgt acagatgcaa gcaccccgc aaatcgacaa cgcacgaaag 2640
198 atccgaaaga gacgattgta gccgataagc cagtcgatat tccgtcatca cgtaccgtag 2700
199 ctccgacact tttgactatg atgggtcacg aagatcagca gaaaacgtgc gctgcaacgc 2760
200 ccggttgttg ttcgtgttcg gctatcgaag aatcatcgga acagaacaag aagaagcgac 2820
201 cggatccgat gtcggcgatc gaatcatctg catttgagaa taagctgttg gatgaggttt 2880
202 taatgccgag agacacgatg cgagtgagac gatcaattga agacgcgaat cgagtcagtg 2940
203 aagagttgga aaaagctgaa aatttgggaa aagctccaaa aactctcggt ggaaagaagc 3000
204 cgctgatcca tatttcgaag aagaagccgt cgagcagcag caccacatcc acaccggctc 3060
205 caacgatcgc atcaatgtat gccttaacaa ggaaaccgac tacggtgccg ggaacaagga 3120
206 ttcggctcta cgagatctac gaacctttac ccggaagctg ggcgattaat gtatcagctc 3180
207 tggcattgga taatagttat gtgatacgaa atttgaagca ttacacactt tatgcgattt 3240
208 ctctatccgc gtgccaaaac atgacagtac ccggagcatc ttgctcaata tcccatcgtg 3300
209 cgggagcatt gaaacgaaca aaacacatca cagacattga taaagtgttg aatgaaacaa 3360
210 ttgaatggag atttatgaat aatagtcaac aagtcaacgt gacgtgggat ccaccgactg 3420
211 aagtgaatgg tggaatattc ggttatgttg taaagcttaa gtcaaaagtc gatggatcaa 3480
212 ttgttatgac gagatgtgtc ggtgcgaaga gaggatattc aacacggaat cagggtgtcc 3540
213 tattccagaa tttggccgat ggacgttatt ttgtctcagt aacggcgacc tctgtacacg 3600
214 gcgctggacc ggaagccgaa tcctccgacc caatcgtcgt catgacgcca ggcttcttca 3660
215 ctgtggaaat cattctcggc atgcttctcg tctttttgat tttaatgtca attgccggtt 3720
216 gtataatcta ctactacatt caagtacgct acggcaaaaa agtgaaagct ctatctgact 3780
217 ttatgcaatt gaatcccgaa tattgtgtgg acaataagta caatgcagac gattgggagc 3840
218 tacggcagga tgatgttgtg ctcggacaac agtgtggaga gggatcattc ggaaaagtgt 3900
219 acctaggaac tggaaataat gttgtttctc tgatgggtga tcgtttcgga ccgtgtgcta 3960
220 ttaagattaa tgtagatgat ccagcgtcga ctgagaatct caactatctc atggaagcta 4020
221 atattatgaa gaactttaag actaactta tcgtccaact gtacggagtt atctctactg 4080
222 tacaaccagc gatggttgtg atggaaatga tggatcttgg aaatctccgt gactatctcc 4140
223 gatcgaaacg cgaagacgaa gtgttcaatg agacggactg caactttttc gacataatcc 4200
224 cgagggataa attccatgag tgggccgcac agatttgtga tggtatggcg tacctggagt 4260
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226 ctgtcaagat tggagatttc ggaatggctc gtgatctatt ctatcatgac tattataagc 4380
227 catcgggcaa gcgtatgatg cctgttcgat ggatgtcacc cgagtcgttg aaagacggaa 4440
228 agtttgactc gaaatctgat gtttggagct cggagttgt tctctatgaa atggttacac 4500
229 tcggtgctca gccatatatt ggtttgagta atgatgaggt gttgaattat attggaatgg 4560
230 cccggaaggt tatcaagaag cccgaatgtt gtgaaaacta ttggtataag gtgatgaaaa 4620
231 tgtgctggag atactcacct cgggatcgtc cgacgttcct ccagctcgtt catcttctag 4680
232 cagctgaagc ttcaccagaa ttccgagatt tatcatttgt cctaaccgat aatcaaatga 4740
233 tccttgacga ttcagaagca ctggatcttg atgatattga tgatactgat atgaatgatc 4800
234 aggttgtcga ggtggcaccg gatgttgaga acgtcgaggt tcagagtgat tcggaacgtc 4860
235 ggaatacgga ttcaataccg ttgaaacagt ttaagacgat ccctccgatc aatgcgacga 4920
236 cgagtcattc gacaatatcg attgatgaga caccgatgaa agcgaagcag cgagaaggat 4980
237 cgctggatga ggagtacgca ttgatgaatc atagtggagg tccgagtgat gcggaagttc 5040
```

```
238 ggacgtatgc tggtgatgga gattatgtgg agagagatgt tcgagagaat gatgtgccaa   5100
239 cgcgacgaaa tactggtgca tcaacatcaa gttacacagg tggtggtcca tattgcctaa   5160
240 caaatcgtgg tggttcaaat gaacgaggag ccggtttcgg tgaagcagta cgattaactg   5220
241 atggtgttgg aagtggacat ttaaatgatg atgattatgt tgaaaaagag atatcatcca   5280
242 tggatacgcg ccggagcacg ggcgcctcga gctcttccta cggtgttcca cagacgaatt   5340
243 ggagtggaaa tcgtggtgcc acgtattata cgagtaaagc tcaacaggca gcaactgcag   5400
244 cagcagcagc agcagcagct ctccaacagc aacaaatgg tggtcgaggc gatcgattaa    5460
245 ctcaactacc cggaactgga catttacaat cgacacgtgg tggacaagat ggagattata   5520
246 ttgaaactga accgaaaaat tatagaaata atggatctcc atcgcgaaac ggcaacagcc   5580
247 gtgacatttt caacggacgt tcggctttcg gtgaaaatga gcatctaatc gaggataatg   5640
248 agcatcatcc acttgtctga aacccccaaa aaatcccgcc tcttaaatta taaattatct   5700
249 cccacattat catatctcta cacgaatatc ggattttttt tcagattttt tctgaaaaat   5760
250 tctgaataat tttaccccat ttttcaaatc tctgtatttt tttttgttat tacccc        5816
254 <210> SEQ ID NO: 12
255 <211> LENGTH: 1724
256 <212> TYPE: PRT
257 <213> ORGANISM: Caenorhabditis elegans
259 <400> SEQUENCE: 12
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261 1               5                   10                  15
262 Ala Ser Ile Val Glu Lys Arg Cys Gly Pro Ile Asp Ile Arg Asn Arg
263             20              25                  30
264 Pro Trp Asp Ile Lys Pro Gln Trp Ser Lys Leu Gly Asp Pro Asn Glu
265         35                  40                  45
266 Lys Asp Leu Ala Gly Gln Arg Met Val Asn Cys Thr Val Val Glu Gly
267     50                  55                  60
268 Ser Leu Thr Ile Ser Phe Val Leu Lys His Lys Thr Lys Ala Gln Glu
269 65                  70                  75                  80
270 Glu Met His Arg Ser Leu Gln Pro Arg Tyr Ser Gln Asp Glu Phe Ile
271             85                  90                  95
272 Thr Phe Pro His Leu Arg Glu Ile Thr Gly Thr Leu Leu Val Phe Glu
273             100                 105                 110
274 Thr Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val
275         115                 120                 125
276 Ile Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg
277     130                 135                 140
278 Asn Pro Asp Leu Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn
279 145                 150                 155                 160
280 Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr
281             165                 170                 175
282 Ile Asp Trp Lys His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val
283             180                 185                 190
284 Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys Pro
285         195                 200                 205
286 Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr Leu
287     210                 215                 220
288 Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys Trp
289 225                 230                 235                 240
290 Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro
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Please Note:
**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.**

Seq#:11; N Pos. 2050
Seq#:31; N Pos. 3,12,15,18,21
Seq#:32; N Pos. 7,8,9,12,15
Seq#:115; Xaa Pos. 4,5,11,12,16,37,38,39,41,42,43,47
Seq#:126; Xaa Pos. 20,21,22
Seq#:127; Xaa Pos. 20,21,22
Seq#:128; Xaa Pos. 20,21,22
Seq#:129; Xaa Pos. 20,21,22
Seq#:130; Xaa Pos. 20,21,22
Seq#:131; Xaa Pos. 20,21,22
Seq#:132; Xaa Pos. 20,21,22
Seq#:133; Xaa Pos. 20,21,22
Seq#:134; Xaa Pos. 20,21,22
Seq#:135; Xaa Pos. 20,21,22
Seq#:136; Xaa Pos. 20,21,22
Seq#:137; Xaa Pos. 20,21,22
Seq#:138; Xaa Pos. 20,21,22
Seq#:139; Xaa Pos. 20,21,22
Seq#:140; Xaa Pos. 20,21,22
Seq#:141; Xaa Pos. 20,21,22
Seq#:142; Xaa Pos. 20,21,22
Seq#:143; Xaa Pos. 20,21,22
Seq#:144; Xaa Pos. 20,21,22
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Seq#:147; Xaa Pos. 20,21,22
Seq#:148; Xaa Pos. 20,21,22
Seq#:149; Xaa Pos. 20,21,22
Seq#:150; Xaa Pos. 20,21,22
Seq#:151; Xaa Pos. 20,21,22
Seq#:152; Xaa Pos. 20,21,22
Seq#:153; Xaa Pos. 20,21,22
Seq#:238; Xaa Pos. 84,85,86,87,88,89,90,91,92,93,94,95,96
Seq#:304; Xaa Pos. 4,5
Seq#:323; Xaa Pos. 2,3,5,6

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L:188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:2040
L:799 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:803 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:31
L:804 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:814 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:818 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:32
L:819 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:3338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115 after pos.:0
M:341 Repeated in SeqNo=115
L:3560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:126 after pos.:16
L:3578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127 after pos.:16
L:3596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:128 after pos.:16
L:3614 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:129 after pos.:16
L:3632 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:130 after pos.:16
L:3650 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131 after pos.:16
L:3668 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132 after pos.:16
L:3686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:133 after pos.:16
L:3704 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:134 after pos.:16
L:3722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:135 after pos.:16
L:3740 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:136 after pos.:16
L:3758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:137 after pos.:16
L:3776 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138 after pos.:16
L:3794 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139 after pos.:16
L:3812 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:140 after pos.:16
L:3830 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:141 after pos.:16
L:3848 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:142 after pos.:16
L:3866 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:143 after pos.:16
L:3884 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:144 after pos.:16
L:3902 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:145 after pos.:16
L:3920 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146 after pos.:16
L:3938 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:147 after pos.:16
L:3956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:148 after pos.:16
L:3974 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:149 after pos.:16
L:3992 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:150 after pos.:16
L:4010 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:151 after pos.:16
L:4028 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:152 after pos.:16
L:4046 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:153 after pos.:16
L:5914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:238 after pos.:80
L:6809 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:304 after pos.:0
L:7349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:323 after pos.:0
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